

## PATENT COOPERATION TREATY

From the INTERNATIONAL BUREAU

PCT

## NOTIFICATION OF ELECTION

(PCT Rule 61.2)

To:

United States Patent and Trademark  
Office  
(Box PCT)  
Crystal Plaza 2  
Washington, DC 20231  
ÉTATS-UNIS D'AMÉRIQUE

in its capacity as elected Office

Date of mailing (day/month/year) 12 March 1999 (12.03.99)	
International application No. PCT/US98/15411	Applicant's or agent's file reference 234/231 PCT
International filing date (day/month/year) 24 July 1998 (24.07.98)	Priority date (day/month/year) 25 July 1997 (25.07.97)
<b>Applicant</b> BRIGGS, Michael, R. et al	

1. The designated Office is hereby notified of its election made:

in the demand filed with the International Preliminary Examining Authority on:

12 February 1999 (12.02.99)

in a notice effecting later election filed with the International Bureau on:

\_\_\_\_\_

2. The election  was

was not

made before the expiration of 19 months from the priority date or, where Rule 32 applies, within the time limit under Rule 32.2(b).

The International Bureau of WIPO 34, chemin des Colombettes 1211 Geneva 20, Switzerland  Facsimile No.: (41-22) 740.14.35	Authorized officer  A. Karkachi  Telephone No.: (41-22) 338.83.38
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## PATENT COOPERATION TREATY

REC'D 21 MAY 1999

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WIPO PCT

19

## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

(PCT Article 36 and Rule 70)

Applicant's or agent's file reference 234/231-PCT	FOR FURTHER ACTION	See Notification of Transmittal of International Preliminary Examination Report (Form PCT/IPEA/416)
International application No. PCT/US98/15411	International filing date (day/month/year) 24 JULY 1998	Priority date (day/month/year) 25 JULY 1997
International Patent Classification (IPC) or national classification and IPC Please See Supplemental Sheet.		
Applicant LIGAND PHARMACEUTICALS INCORPORATED		

1. This international preliminary examination report has been prepared by this International Preliminary Examining Authority and is transmitted to the applicant according to Article 36.

2. This REPORT consists of a total of 5 sheets.

This report is also accompanied by ANNEXES, i.e., sheets of the description, claims and/or drawings which have been amended and are the basis for this report and/or sheets containing rectifications made before this Authority. (see Rule 70.16 and Section 607 of the Administrative Instructions under the PCT).

These annexes consist of a total of 0 sheets.

3. This report contains indications relating to the following items:

- I  Basis of the report
- II  Priority
- III  Non-establishment of report with regard to novelty, inventive step or industrial applicability
- IV  Lack of unity of invention
- V  Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement
- VI  Certain documents cited
- VII  Certain defects in the international application
- VIII  Certain observations on the international application

Date of submission of the demand 12 FEBRUARY 1999	Date of completion of this report 06 MAY 1999
Name and mailing address of the IPEA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Facsimile No. (703) 305-3230	Authorized officer IREM YUCEL JOYCE BRIDGERS PARALEGAL SPECIALIST CHEMICAL MATRIX Telephone No. (703) 308-0196 <i>JMB</i>

## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No.  
PCT/US98/15411**I. Basis of the report**

1. This report has been drawn on the basis of (*Substitute sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to the report since they do not contain amendments*):

- the international application as originally filed.
- the description, pages 1-86, as originally filed.  
pages NONE, filed with the demand.  
pages NONE, filed with the letter of \_\_\_\_\_.  
pages \_\_\_\_\_, filed with the letter of \_\_\_\_\_.
- the claims, Nos. 1-25, as originally filed.  
Nos. NONE, as amended under Article 19.  
Nos. NONE, filed with the demand.  
Nos. NONE, filed with the letter of \_\_\_\_\_.  
Nos. \_\_\_\_\_, filed with the letter of \_\_\_\_\_.
- the drawings, sheets/fig 1-7, as originally filed.  
sheets/fig NONE, filed with the demand.  
sheets/fig NONE, filed with the letter of \_\_\_\_\_.  
sheets/fig \_\_\_\_\_, filed with the letter of \_\_\_\_\_.

2. The amendments have resulted in the cancellation of:

- the description, pages none.
- the claims, Nos. none.
- the drawings, sheets/fig none.

3.  This report has been established as if (some of) the amendments had not been made, since they have been considered to go beyond the disclosure as filed, as indicated in the **Supplemental Box**. Additional observations below (Rule 70.2(c)).

4. Additional observations, if necessary:

NONE

**III. Non-establishment of opinion with regard to novelty, inventive step and industrial applicability**

The question whether the claimed invention appears to be novel, to involve an inventive step (to be non-obvious), or to be industrially applicable have not been and will not be examined in respect of:

- the entire international application.
- claims Nos. 9-14.

because:

- the said international application, or the said claim Nos.    relate to the following subject matter which does not require international preliminary examination (*specify*).

- the description, claims or drawings (*indicate particular elements below*) or said claims Nos.    are so unclear that no meaningful opinion could be formed (*specify*).

- the claims, or said claims Nos.    are so inadequately supported by the description that no meaningful opinion could be formed.

- no international search report has been established for said claims Nos. 9-14.

## INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No.  
PCT/US98/15411

**V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement****1. STATEMENT**

Novelty (N)	Claims <u>1-8 and 15-25</u>	YES
	Claims <u>none</u>	NO
Inventive Step (IS)	Claims <u>1-8 and 15-25</u>	YES
	Claims <u>none</u>	NO
Industrial Applicability (IA)	Claims <u>1-8 and 15-25</u>	YES
	Claims <u>none</u>	NO

**2. CITATIONS AND EXPLANATIONS**

Claims 1-8 and 15-25 meet the criteria set out in PCT Article 33(2)-(4), because the prior art does not teach or fairly suggest the control regions of the human PPAR $\gamma$  gene. The prior art does not teach nor fairly suggest methods of screening for agents which modulate expression of the human PPAR $\gamma$  gene by using the control region of the human PPAR $\gamma$  gene in reporter gene constructs. Further the claimed invention recited by the instant claims has industrial applicability.

----- NEW CITATIONS -----  
NONE

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No.

PCT/US98/15411

**Supplemental Box**

(To be used when the space in any of the preceding boxes is not sufficient)

Continuation of: Boxes I - VIII

Sheet 10

**CLASSIFICATION:**

The International Patent Classification (IPC) and/or the National classification are as listed below:

IPC(6): C07H 21/02, 21/04; C12N 5/06, 5/08, 5/10; C12Q 1/02, 1/68; C12P 21/06 and US Cl.: 536/23.1, 23.5, 24.1; 435/6, 29, 69.1, 357, 365, 370

## INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US98/15411

**A. CLASSIFICATION OF SUBJECT MATTER**

IPC(6) :Please See Extra Sheet.

US CL :536/23.1, 23.5, 24.1; 435/6, 29, 69.1, 357, 365, 370

According to International Patent Classification (IPC) or to both national classification and IPC

**B. FIELDS SEARCHED**

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 536/23.1, 23.5, 24.1; 435/6, 29, 69.1, 357, 365, 370

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

NONE

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

Please See Extra Sheet.

**C. DOCUMENTS CONSIDERED TO BE RELEVANT**

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A, P	US 5,686,596 A (MUKHERJEE) 11 November 1997, see entire document.	1-8, 15-25
A, P	FAJAS et al. The Organization, Promoter Analysis, and Expression of the Human PPAR gamma Gene. The Journal of Biological Chemistry. 25 July 1997, Vol. 272, No. 30, pages 18799-18789, see entire document.	1-8, 15-25
Y, P	US 5,726,041 A (CHRESPI et al.) 10 March 1998, see entire document.	1-8, 15-25
Y	GEARING et al. Structure of the Mouse Peroxisome Proliferator Activated Receptor alpha Gene. Biochemical and Biophysical Research Communications. 28 February 1994, Vol. 199, No. 1, pages 255-263, see entire document.	1-8, 15-25

Further documents are listed in the continuation of Box C.  See patent family annex.

* Special categories of cited documents:	"T"	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"A" document defining the general state of the art which is not considered to be of particular relevance	"X"	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"B" earlier document published on or after the international filing date	"Y"	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reasons (as specified)	"&"	document member of the same patent family
"O" document referring to an oral disclosure, use, exhibition or other means		
"P" document published prior to the international filing date but later than the priority date claimed		

Date of the actual completion of the international search

25 SEPTEMBER 1998

Date of mailing of the international search report

28 OCT 1998

Name and mailing address of the ISA/US  
Commissioner of Patents and Trademarks  
Box PCT  
Washington, D.C. 20231

Facsimile No. (703) 305-3230

Authorized officer  
*Dorothy Lawrence Jr.*  
IREM YUCEL

Telephone No. (703) 308-0196

## INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US98/15411

## C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	ZHU et al. Structural organization of mouse peroxisome proliferator-activated receptor gamma (mPPARgamma) gene: Alternative promoter use and different splicing yield two mPPR gamma isoforms. Proceedings of National Academy of Sciences, U.S.A. August 1995, Vol. 92, pages 7921-7925, see entire document.	1-8, 15-25

**INTERNATIONAL SEARCH REPORT**International application No.  
PCT/US98/15411**Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)**

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1.  Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
  
2.  Claims Nos.: 9-14 because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:  

Please See Extra Sheet.
  
3.  Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

**Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)**

This International Searching Authority found multiple inventions in this international application, as follows:

1.  As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
  
2.  As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
  
3.  As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
  
4.  No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

**Remark on Protest**  

- The additional search fees were accompanied by the applicant's protest.  
No protest accompanied the payment of additional search fees.

**INTERNATIONAL SEARCH REPORT**

International application No.  
PCT/US98/15411

**A. CLASSIFICATION OF SUBJECT MATTER:**

IPC (6):

C07H 21/02, 21/04; C12N 5/06, 5/08, 5/10; C12Q 1/02, 1/68; C12P 21/06

**B. FIELDS SEARCHED**

Electronic data bases consulted (Name of data base and where practicable terms used):

APS, STN (CAPLUS), DIALOG (MEDLINE, BIOSIS, SCISEARCH)

TERMS: PPAR, peroxisome proliferator activated receptor?, genom? clon? untranslate? region? sequence? regulat? control? element? human?

**BOX I. OBSERVATIONS WHERE CLAIMS WERE FOUND UNSEARCHABLE**

## 2. Where no meaningful search could be carried out, specifically:

The above claims are drawn to specific sequences or specific regions (subsequences) of particular genes. The CRF submitted in response to a telephone call to Applicant's representative did not comply with the sequence rules. Applicant's representative was FAXed a copy of the error report, but has not yet submitted a subsequent CRF.

***FOR THE PURPOSES OF INFORMATION ONLY***

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

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AZ	Azerbaijan	GB	United Kingdom	MC	Monaco	TD	Chad
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BR	Brazil	IL	Israel	MW	Malawi	UG	Uganda
BY	Belarus	IS	Iceland	MX	Mexico	US	United States of America
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CN	China	KZ	Kazakhstan	RO	Romania		
CU	Cuba	LC	Saint Lucia	RU	Russian Federation		
CZ	Czech Republic	LI	Liechtenstein	SD	Sudan		
DE	Germany	LK	Sri Lanka	SE	Sweden		
DK	Denmark	LR	Liberia	SG	Singapore		
EE	Estonia						

1

## SEQUENCE LISTING

<110> Michael R. Briggs  
Regis S. Saladin  
Johan Auwerx  
Lluis Fajas

<120> HUMAN PEROXISOME PROLIFERATOR ACTIVATED RECEPTOR GAMMA  
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<130> 234/231-PCT

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<213> Human PPAR $\gamma$ 1 promoter

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<213> Human PPAR $\gamma$ 2 promoter, exon B, and intron B

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&lt;211&gt; 30

&lt;212&gt; DNA

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&lt;210&gt; 9

&lt;211&gt; 29

5

&lt;212&gt; DNA

&lt;213&gt; LF-22

&lt;400&gt; 9

gcaattgaat gtcgtgtctg tggagataa

29

&lt;210&gt; 10

&lt;211&gt; 29

&lt;212&gt; DNA

&lt;213&gt; LF-23

&lt;400&gt; 10

gtggatccga cagttaaat cacatctgt

29

&lt;210&gt; 11

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; LF-24

&lt;400&gt; 11

gttagaaataa atgtcagttac tgcgggttcc

30

&lt;210&gt; 12

&lt;211&gt; 29

&lt;212&gt; DNA

&lt;213&gt; LF-25

&lt;400&gt; 12

tcgatatacac tggagatctc cgccaaacag

29

&lt;210&gt; 13

&lt;211&gt; 30

6

&lt;212&gt; DNA

&lt;213&gt; LF-26

&lt;400&gt; 13

acataaaagtc cttcccgctg accaaagcaa

30

&lt;210&gt; 14

&lt;211&gt; 29

&lt;212&gt; DNA

&lt;213&gt; LF-27

&lt;400&gt; 14

ctctgctcct gcaggggggt gatgtgttt

29

&lt;210&gt; 15

&lt;211&gt; 29

&lt;212&gt; DNA

&lt;213&gt; LF-28

&lt;400&gt; 15

gaagttcaat gcactggaat tagatgaca

29

&lt;210&gt; 16

&lt;211&gt; 29

&lt;212&gt; DNA

&lt;213&gt; LF-29

&lt;400&gt; 16

gagctccagg ggtagtagca gggtgtctt

29

&lt;210&gt; 17

&lt;211&gt; 28

&lt;212&gt; DNA

7

&lt;213&gt; LF-33

&lt;400&gt; 17

gacgggctga ggagaagtca cactctga

28

&lt;210&gt; 18

&lt;211&gt; 28

&lt;212&gt; DNA

&lt;213&gt; LF-35

&lt;400&gt; 18

agcatggaat aggggtttgc tgtaattc

28

&lt;210&gt; 19

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; LF-36

&lt;400&gt; 19

tagtacaagt ccttgttagat ctcc

24

&lt;210&gt; 20

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; LF-44

&lt;400&gt; 20

gtcggcctcg aggacaccgg agag

24

&lt;210&gt; 21

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; LF-58

&lt;400&gt; 21

caactcatgtg acaagacctg ctcc

24

&lt;210&gt; 22

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; LF-59

&lt;400&gt; 22

gccgacacta aaccaccaat atac

24

&lt;210&gt; 23

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; LF-60

&lt;400&gt; 23

cgttaaaggc tgactctcggt ttga

24

&lt;210&gt; 24

&lt;211&gt; 26

&lt;212&gt; DNA

&lt;213&gt; AII J PPRE

&lt;400&gt; 24

gatccttcaa cctttaccct ggtaga

26

&lt;210&gt; 25

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; ACO PPRE

9

&lt;400&gt; 25

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30

&lt;210&gt; 26

&lt;211&gt; 27

&lt;212&gt; DNA

&lt;213&gt; LPL PPRE

&lt;400&gt; 26

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27

&lt;210&gt; 27

&lt;211&gt; 19

&lt;212&gt; DNA

<213>  $\gamma$  AS

&lt;400&gt; 27

gcattatgag catccccac

19

&lt;210&gt; 28

&lt;211&gt; 20

&lt;212&gt; DNA

<213>  $\gamma$ S

&lt;400&gt; 28

tctctccgtatggaaagacc

20

&lt;210&gt; 29

&lt;211&gt; 19

&lt;212&gt; DNA

<213>  $\gamma$ 2S

10

&lt;400&gt; 29

gcgattcctt cactgatac

19

&lt;210&gt; 30

&lt;211&gt; 52

&lt;212&gt; DNA

&lt;213&gt; CDS

&lt;400&gt; 30

ttctagaatt cagcgccgc tttttttttt tttttttttt tttttttttt vn

52

&lt;210&gt; 31

&lt;211&gt; 201

&lt;212&gt; DNA

<213> PPAR $\gamma$ 1 proximal promoter

&lt;400&gt; 31

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 ccgaccccgga tccgcccggc cgggcaggcg gggcccagcg cactcgagc ccgagcccgaa  
 gccgcagcccg ccgcctgggg cgcttgggtc ggcctcgagg acaccggaga ggggcgcac  
 gccgcccgtgg ccgcagaaat g

60

120

180

201

&lt;210&gt; 32

&lt;211&gt; 177

&lt;212&gt; DNA

<213> PPAR $\gamma$ 2 proximal promoter

&lt;400&gt; 32

 gtcctttctg tgtttattcc catctctccc aaatatgg aaactgatgt ctgtactcat  
 gggtgtattc acgattctgt tacttcaagt cttttcttt taacggattg atcttttgct  
 agatagagac aaaatatcag tgtgaattac agcaaaccac tattccatgc tgttatg

60

120

177

&lt;210&gt; 33

&lt;211&gt; 468

&lt;212&gt; DNA

<213> PPAR $\gamma$ 3 proximal promoter

&lt;400&gt; 33

taatcctttt aaggcttagt ttttcttaag tctgtgcagt aatagaggta tcgtcattca	60
tgtgacataa aagatggaaa ggggcttcat tcatagttagt gatggaaata ggaaaggtagg	120
tgaagtgatt ttaatagatg ttcttttat gaaataattt ttaaagattt tccagccctg	180
catgatttat gatgaatcat ttgtgtgtct gttagttact tttagagaat agaaagcatt	240
gtaggcttag ggaagcaaa cattcagaat gaaatccaat agagaaggta aatttatttg	300
ggcatgtaca ttggcagc ctaggctgt tacatgtgtcacattctga acatgtgtgt	360
atattgaaaa tcttgtctct tttttattgt taagatttga aagaagccga cactaaacca	420
ccaatataca acaaggccat ttgtcaaacc gagagtcaac cttaacg	468

&lt;210&gt; 34

&lt;211&gt; 1433

&lt;212&gt; DNA

<213> PPAR $\gamma$ 3 promoter, exon A2, and intron A2

&lt;400&gt; 34

gagaatacag gcacatgcca ccatgcccag ctaatttttc tggtttttgt agagacagga	60
tttcgctgtg gtgctcaggc tggctccaa ctcctggct caagcaatcc gcctgcctca	120
gccttccaaa gtgaaaagggt ttctctcat ttctcaaata gaagtactaa acaatgccag	180
agaaataaaat aaacaggcaa aatacgttgg ctatagtttta tattatttcc tgctacagtt	240
aacaaaatgg gaagacattt tatcttcatg gtctactaca ttatgccat gtgttaagta	300
ataaaaatagc ttttgtaaat tataaattaa aagtcagata tttaaaagag aaaatactgt	360
agagtttca tggtagttaag actgtgtaga atgtcggtc tggatgttgg cgctattcaa	420
gcctgtatgtc taaggctttt ggcatttagat gctgtttgt ctcatggaa aatacagcta	480
ttcttaggatc ttggagccct tcataagaga taagggtgtg aatcctaaga ccctaggacc	540
atttacttag atgatctgtc ctctgggtcg tcctctgaaa agtctgttc gtgaggggtg	600
tgctgcattt gccttccta agtgggtgtgg cacacaactg tactgtcacc tttaggcttaa	660
taaccatgtg tcatactagaa tgaagttata tttaaaaag gatcggtttt gccatgtata	720
aattttcaaa cattaacttt cagggttatt aatccttttta aggtctagtt ttcttaagt	780
ctgtgcagta atagaggat cgtcattcat gtgacataaa agatggaaag gggcttcatt	840
catgttagtg atggaaaatag gaaagttagt gaagtgtttttaatagatgt ttctttatgt	900
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taagatttga aagaagccga cactaaacca ccaatataca acaaggccat ttctcaaacc	1200
gagagtcaac ctttaacggc aagtaaaatc agaatttata ctgcattgtt attgaaaagt	1260
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gtgatttttg actatttata tactttctgc tatataattt tccagtcgt tggactatgc	1380
agtgtaaacat atttgtctaa cacaacaa aggtaaagata ggaaatgac cttagaagttg	1440
agaaataact caaatccctta aaa	1433

<210> 35

<211> 695

<212> DNA

<213> Intron B, exon 1, and intron 1

<400> 35

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tctattatac tcaataaaagc tggacaaaat tttaaataaaa taacagcagt cattaacaga 120  
ctcaattgtat gacctaattgt agaagttaat gagagcaggc ctgttggcaa aaaggcattt 180  
atatggatac actgttatgtt tctgcactgt ttcaaggatcc tctattatga tacctgggta 240  
aagggtgact tcctttctat cataaaacag cctagacagc actaagaagg tggttatgtt 300  
ctttctgtt gttgtgagcg cccagatgag attactttgc caaagactct tttcatttct 360  
ctttctgaaa ctctgtgaga ttgctgtgtt ctctaggact taacttcaca gctagtctat 420  
tttcctttc agaaatgacc atgggtgaca cagagatgcc attctggccc accaactttg 480  
ggatcagctc cgtggatctc tccgtaatgg aagaccactc ccactcctt gatatcaagc 540  
ccttcactac tgttgacttc tccagcattt ctactccaca ttacgaagac attccattca 600  
caagaacaga tccagtggtt gcagattaca agtatgacct gaaacttcaa gagtaccaaa 660  
gtatgatgtt tgtttcaact tttcagacta ctagg 695

<210> 36

<211> 313

<212> DNA

<213> Intron 1, exon 2, and intron 2

<400> 36

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gcctgcattc tccacccattt tattctgaga agactcagct ctacaataag cctcatgaag 120  
gccttc当地 ctccctcatg gcaattgaat gtcgtgtctg tggagataaa gcttctggat 180  
ttcactatgg agttcatgct tgtgaaggat gcaaggtaat taaaaaaaaa gtcttcaaag 240  
aaatttgtga aactttatta tttcatttca gcagaacccc ttttttaggt gatacaatat 300  
atgaattttt ttt 313

<210> 37

<211> 473

<212> DNA

<213> Intron 2, exon 3, and intron 3

<400> 37

gatacccttc gctgttagtt cgtgcttcca tgtgtcataa agactaaaaa tttgcttctt 60

13

tttatccct ttgcaggggtt tcttcggag aacaatcaga ttgaagctta tctatgacag 120  
 atgtatctt aactgtcgga tccacaaaaa aagtagaaat aaatgtcagt actgtcggtt 180  
 tcagaaatgc cttgcagtgg gnatgtctca taatggtaag taaacagtca tcaccatata 240  
 ctttattatt ctcatatag ctgccagacc agtggacact aaagccattg caaaaaatgt 300  
 gtacagtttt tccaccaaattt gccagaattt agaatattgc atggcgataa aacatttctc 360  
 ttttaggtca gtgttttaa agtttatta tagaaccttt ctctctgtgg ttgggcatct 420  
 gccatgagga gaaaagagac ttgaaaaatc tgggggatta tgggaaaaac ctt 473

&lt;210&gt; 38

&lt;211&gt; 706

&lt;212&gt; DNA

&lt;213&gt; Intron 3, exon 4, and intron 4

&lt;400&gt; 38

acaactttga attctgcaca gtttcgtatt ttaattcgtg aaacgtgttg atccttctaa 60  
 gtgcctgacc ttaggtcaag tgctggggat acaaagaagg tgaccttga attgggtctt 120  
 gagggatgag tagaggtgg ttctcaatta tttcacgttt aagtcgacat acttccctcc 180  
 ctttgcataa ctcgaattct ttcactttct cagcaggagt atgcattaac tttaaaaat 240  
 gaaagttAAC ggttaattt ttactgtatgg tctgtctac ttttgtaaaaa taaaaacatg 300  
 agcaaagtgg tagacagaaa ccaggactca agagcagtgg aggaggaggg cttctactgt 360  
 gtgggaacga gggctggag accacagtgt gtgttcagag cagtagtaat ccaatgattc 420  
 atcctgtcat tcctttccct ctatagccat caggtttggg cgatgccac aggccgagaa 480  
 ggagaagctg ttggcggaga tctccagtgat tatcgaccag ctgaatccag agtccgctga 540  
 cctccgtgcc ctggcaaaaac atttgtatga ctcatacata aagtccctcc cgctgaccaa 600  
 agcaaaggcg agggcgatct tgacagggaaa gacaacagac aaatcgtta gttctttct 660  
 gctgtcttca ttgggggagg cggaaagggtt tttgggatt tttgtt 706

&lt;210&gt; 39

&lt;211&gt; 732

&lt;212&gt; DNA

&lt;213&gt; Intron 4, exon 5, and intron 5

&lt;400&gt; 39

ggaaaagaag accaaaattt gtgaaatatg tttggtccca gaagataatt aagatgaata 60  
 aaagaacttg agagtatTTT ctcattatta agcatctca gctttaaaga ttttagttag 120  
 caaagcaagt ttacataaac agtttctga acctgggatg gcattcactg tgagttagaa 180  
 atctccaagt catcccacgt tttccctgtt ttatttgcaag ccattcgtt aatgtacat 240  
 gaattccta atgatggag aagataaaat caagttcaaa cacatcaccc ccctgcagga 300  
 gcagagcaaa gaggtggcca tccgcattt tcagggctgc cagttcgct ccgtggaggg 360  
 tttgcaggag atcacagagt atgccaAAAG cattcctggt tttgtaaatc ttgacttgaa 420  
 cgaccaagta actctcctca aatatggagt ccacgagato atttacacaa tgctggccctc 480  
 cttgatgaat aaagatgggg ttctcatatc cgagggccaa ggcttcatga caagggagtt 540  
 tctaaagagc ctgcgaaagc cttttggta ctttatggag cccaagttt agtttgcgtt 600  
 gaagttcaat gcactggaat tagatgacag cgacttggca atatttattt ctgtcattat 660  
 ttcagtggta ggttaagattt gtcttttgc tttctatgaa agagggtggg atgatgggg 720  
 ggtggccaaa ag 732

14

<210> 40  
<211> 592  
<212> DNA  
<213> Intron 5, exon 6, and 3' UTR  
<400> 40

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gagntgcnta ggcctccaag gcggggccca gaggatttt tgactgaacc ccctgttgt 120  
tttccatat gtgctcccc agaccgccc ggttgctga atgtgaagcc cattgaagac 180  
attcaagaca acctgctaca agccctggag ctccagctga agctgaacca ccctgagtcc 240  
tcacagctgt ttgccaagct gctccagaaa atgacagacc tcagacagat tgcacggaa 300  
cacgtgcagc tactgcaggt gatcaagaag acggagacag acatgagtct tcacccgctc 360  
ctgcaggaga tctacaagga cttgtactag cagagagtcc tgagccactg ccaacatttc 420  
ccttcttcca gttgcactat tctgagggaa aatctgacca taagaaattt actgtgaaaa 480  
agcgttttaa aaagaaaagg gtttagata tgcattttt tatgcattt gttataaag 540  
acacattttac aatttacttt taatattaaa aattaccata ttatgaaattt gc 592

<210> 41  
<211> 13  
<212> DNA  
<213> PPAR $\gamma$ 3-E-box  
<400> 41

attcatgtga cat

13

<210> 42  
<211> 13  
<212> DNA  
<213> PPAR $\gamma$ 3-E-box  
<400> 42

attcatgcat cat

13

<210> 43  
<211> 13

15

&lt;212&gt; DNA

&lt;213&gt; A1 (97) Donor

&lt;400&gt; 43

cgcaggtaat agt

13

&lt;210&gt; 44

&lt;211&gt; 13

&lt;212&gt; DNA

&lt;213&gt; A1 (97) Acceptor

&lt;400&gt; 44

ttgttaagat ttg

13

&lt;210&gt; 45

&lt;211&gt; 13

&lt;212&gt; DNA

&lt;213&gt; A2 (74) Donor

&lt;400&gt; 45

taacggtaag taa

13

&lt;210&gt; 46

&lt;211&gt; 13

&lt;212&gt; DNA

&lt;213&gt; A2 (74) Acceptor

&lt;400&gt; 46

cctttcagaa atg

13

&lt;210&gt; 47

&lt;211&gt; 12

16

&lt;212&gt; DNA

&lt;213&gt; B (211) Donor

&lt;400&gt; 47

caaggtaaag tt

12

&lt;210&gt; 48

&lt;211&gt; 13

&lt;212&gt; DNA

&lt;213&gt; B (211) Acceptor

&lt;400&gt; 48

cctttcagaa atg

13

&lt;210&gt; 49

&lt;211&gt; 12

&lt;212&gt; DNA

&lt;213&gt; 1 (213) Donor

&lt;400&gt; 49

caaagtatga tg

12

&lt;210&gt; 50

&lt;211&gt; 13

&lt;212&gt; DNA

&lt;213&gt; 1 (231) Acceptor

&lt;400&gt; 50

atacacacaggc gca

13

&lt;210&gt; 51

&lt;211&gt; 12

17

&lt;212&gt; DNA

&lt;213&gt; 2 (170) Donor

&lt;400&gt; 51

caaggttaatt aa

12

&lt;210&gt; 52

&lt;211&gt; 12

&lt;212&gt; DNA

&lt;213&gt; 2 (170) Acceptor

&lt;400&gt; 52

ctttgcaggg tt

12

&lt;210&gt; 53

&lt;211&gt; 12

&lt;212&gt; DNA

&lt;213&gt; 3 (139) Donor

&lt;400&gt; 53

aatggtaagt aa

12

&lt;210&gt; 54

&lt;211&gt; 13

&lt;212&gt; DNA

&lt;213&gt; 3 (139) Acceptor

&lt;400&gt; 54

ctctatagcc atc

13

&lt;210&gt; 55

&lt;211&gt; 12

18

&lt;212&gt; DNA

&lt;213&gt; 4 (203) Donor

&lt;400&gt; 55

atcagttagt tc

12

&lt;210&gt; 56

&lt;211&gt; 12

&lt;212&gt; DNA

&lt;213&gt; 4 (203) Acceptor

&lt;400&gt; 56

atttgcagcc at

12

&lt;210&gt; 57

&lt;211&gt; 12

&lt;212&gt; DNA

&lt;213&gt; 5 (451) Donor

&lt;400&gt; 57

ggaggtaaga tt

12

&lt;210&gt; 58

&lt;211&gt; 13

&lt;212&gt; DNA

&lt;213&gt; 5 (451) Acceptor

&lt;400&gt; 58

ttccccagac cgc

13

&lt;210&gt; 59

&lt;211&gt; 12

19

&lt;212&gt; DNA

&lt;213&gt; 6 (248) Donor

&lt;400&gt; 59

tactagcaga ga

12

&lt;210&gt; 60

&lt;211&gt; 44

&lt;212&gt; DNA

&lt;213&gt; Adaptor

&lt;400&gt; 60

ctaatacgac tcactatagg gctcgagcgg ccgccccggc aggt